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Post-processing: Minimum Match 0%
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Listing first 45 summaries
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Pred. No. is the score greater that and is derived is the number of results predicted by chance to have a ater than or equal to the score of the result being printed, rived by analysis of the total score distribution.

Novel human polynu	AAF66485	22		11.0	371.2	11	
Nucleotide sequenc	AAA64601	21		11.2	376.4	10	
Probe #22725 used	AAI54039	22		13.5	455.2	c 9	
Probe #9583 used t	AA140897	22		15.1	507.2	с В	
Human colon cancer	AAH33745	22		17.2	577.2	7	
Interferon induced	AAS01152	22		19.7	663	6	
Interferon induced	AAS01150	22		19.7	663	5	
Interferon induced	AAS01151	22		23.9	803	4	
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AAH03175	AAH18139	AAQ90049	AAQ76554	AAF58255	AAF58262	AAF58259	AAF58257	AAF58254	AAF58252	AAI23760	AAF58255	AAF58262	AAF58259	AAF58257	AAF58254	AAF58252	AAC77190	AAI24291	AAI14562	AAH15201	AA115107	AA154310	AAI41271	AAI00596	AAI31919	AAI10666	AAQ76640	AAI05627	AAI45108	AAI19913	AAQ90050	AAF66499	AAV89625
Human cDNA clone (Human cDNA sequenc	Melanoma different	Human genome fragm	Oligonucleotide D1	Oligonucleotide D2		Oligonucleotide D1	Oligonucleotide Dl	Oligonucleotide D1	Probe #13693 for g			Oligonucleotide D2	Oligonucleotide Dl	Oligonucleotide D1	Oligonucleotide D1	Human ORFX ORF2745	Probe #14224 for g	Probe #4495 for ge	Human cDNA sequenc	Probe #5040 for ge	Probe #22996 used	Probe #9957 used t	Probe #587 used to	Probe #605 used to		Human genome fragm	Probe #5618 used t	Probe #13794 used	Probe #9846 for ge	Melanoma different	Novel human polynu	EST clone CO522.

ALIGNMENTS

RESULT AAI59285 AAI59285; AAI59285 standard; cDNA; 1557 _

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Human polynucleotide SEQ ID NO 1488.

22-OCT-2001 (first entry)

Human; nootropic; immunosuppressant; cytostatic; gene therapy; cancer; peripheral nervous system; neuropathy; central nervous system; CNS; Alzheimer's; Parkinson's disease; Huntington's disease; haemostatic; amyotrophic lateral sclerosis; Shy-Drager Syndrome; chemotactic; chemokinetic; thrombolytic; drug screening; arthritis; inflammation; ieukaemia; ss

Homo sapiens.

WO200153312-A1

26-JUL-2001.

26-DEC-2000; 2000WO-US34263.

21-JAN-2000; 25-APR-2000; 09-JUL-2000; 19-JUL-2000; 03-AUG-2000; 14-SEP-2000; 19-OCT-2000; 29-NOV-2000; 2000US-0488725. 2000US-055317. 2000US-0558042. 2000US-0598042. 2000US-0653131. 2000US-0653450. 2000US-0663191. 2000US-0693036. 2000US-0693036.

(HYSE-) HYSEQ INC

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Best Local Similarity
Matches 1476; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           in gene therapy. A composition containing a polypeptide or polynucleotide of the invention may be used to treat diseases of the peripheral nervous system, such as peripheral nervous injuries, peripheral neuropathy and localised neuropathies and central nervous system disease, such as Alzheimer's, Parkinson's disease, Huntington's disease, amyotrophic lateral sclerosis, and Shy-Drager Syndrome. Other uses include the utilisation of the activities such as: Inmune system suppression, activing in activity, chemotactic/chemokinetic activity, haemostatic and thrombolytic activity, cancer diagnosis and therapy, drug screening, assays for receptor activity, arthritis and inflammation, leukaemias and
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                      atatgcgctttcccagtggattactgaaaatgaaaaatttgctgaagtaggagtcaaagc
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DB; AAM40129.
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The sequence data for this patent did
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Matches 1355;
                                                                                                                                                                                               system, such as peripheral nervous injuries, peripheral neuropathy and localised neuropathies and central nervous system disease, such as Alzheiner's, Parkinson's disease, Huntington's disease, amyotrophic lateral sclerosis, and Shy-Drager Syndrome. Other uses include the utilisation of the activities such as: Immune system suppression, ActivinyInhibin activity, chemotactic/Chemokinetic activity, haemostatic and thrombolytic activity, cancer diagnosis and therapy, drug screening, assays for receptor activity, arthritis and inflammation, leukaemias and assays for receptor activity, arthritis and inflammation, leukaemias and
                                                                                                                                                                                                                                                                                      The invention relates to human nucleic acids (AAI57798-AAI61369) and the encoded polypeptides (AAM38642-AAM42213) with nootropic, immunosuppressant and cytostatic activity. The polypucleotides are useful in gene therapy. A composition containing a polypeptide or polynucleotide in gene therapy be used to treat diseases of the peripheral nervous of the invention may be used to treat diseases of the peripheral nervous
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as central nervous
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nucleic acids and polypeptides are useful for treating or preventing a pathology associated with IrN polypeptide in a human. They are useful for determining the presence of or predisposition to a disease associated with altered levels of IFN polypeptide or polynucleotide. IFN nucleic acids, polypeptides and antibodies are useful for diagnosis, prevention or treatment of variety of immunological and cell proliferative disorders, such as autoimmune diseases e.g. lupus crythematosus, immunodeficiency diseases such as acquired immunodeficiency syndrome
                                                                                                                                                                                            New interferon induced polypeptides and diagnosis, prevention and treatment of i proliferative disorders, such as lupus each allocations.
                                                                                                                                                                                                                                                                                                                                     (CURA·)
(BIOJ )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Interferon induced nucleic acid; autoimmune disease; lupus erythematosus; immunodeficiency; stroke; acquired immunodeficiency syndrome; AIDS; graft rejection; viral infection; hepatitis; aplastic anaemia; cancer; human immunodeficiency virus; HIV; immune-mediated glomerulonephritis; haematologic disease; chronic neutropenia; myocardial infarction; neurological disease; Alzheimer's disease; parkinson's disease; tumour; meurological disease; Alzheimer's disease; parkinson's disease; tumour;
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20-OCT-1999;
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                                                                                                                                                                                    Alzheimer's disease
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     identifying interferon-like proteins and interferon agonists, for screening drugs and compounds which inhibit or enhance IFN activity or function and as targets for the identification of small molecules that are immunostimulatory, immunosuppressive, or stimulate or suppress normal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            infarction, stroke, neurological diseases including Alzheimer's and Parkinson's diseases, amyotrophic lateral sclerosis and spinal muscular atrophy. IFN nucleic acids and polypeptides are also useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     and cancer. In addition they are also useful for treating or diagnosing various disorders associated with cell death, including myocardial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               immunodeficiency virus (HIV), immune mediated glomerulonephritis, haematologic diseases such as aplastic anaemia and chronic neutro
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       The sequence represents interferon induced nucleic acid, IFN6. IFN nucleic acids and polypeptides are useful for treating or preventing a pathology associated with IFN polypeptide in a human. They are useful for determining the presence of or predisposition to a disease associated with altered levels of IFN polypeptide or polynucleotide. IFN nucleic acids, polypeptides and antibodies are useful for diagnosis, prevention or treatment of variety of immunological and cell proliferative of isomorphism such as autoimmune diseases e.g. lupus erythematosus, immunodeficiency diseases such as acquired immunodeficiency syndrome (AIDS), graft rejection, viral infections including hepatitis and human hammanodeficiency virus (HIV), immune-mediated glomerulonephritis,
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20-OCT-1999;
20-JAN-2000;
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                                                                                                                                                                                              Claim 9; Page 40-43; 134pp; English
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RESULT
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 nucleic acids and polypeptides are useful for treating or preventing a pathology associated with IFN polypeptide in a human. They are useful for determining the presence of or prevention to a disease associated with altered levels of IFN polypeptide or polypucleotide. IFN nucleic acids, polypeptides and antibodies are useful for diagnosis, prevention or treatment of variety of immunological and cell proliferative disorders, such as autoimmune diseases e.g. lupus erythematosus, immunodeficiency diseases such as acquired immunodeficiency syndrome (AIDS), graft rejection, viral infections including hepatitis and human immunodeficiency virus (HIV), immune-mediated glomerulonephritis, heematologic diseases such as aplastic anaemia and chronic neutropenia and cancer. In addition they are also useful for treating or diagnosing various disorders associated with cell death, including mycoardial
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20-OCT-1999;
20-JAN-2000;
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                                                                                                                                                                                                                       The sequence represents interferon induced nucleic acid, IFN5. IFN
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RESULT AAS01152

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Pred. No. 3.3e-154;
0; Mismatches 15;
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or treatment of variety of immunological and cell proliferative disorders, such as autoimmune diseases e.g. lupus erythematosus, immunodeficiency diseases such as acquired immunodeficiency syndrome (AIDS), graft rejection, viral infections including hepatitis and human immunodeficiency virus (HIV), immune-mediated glomerulonephritis, chaematologic diseases such as aplastic anaemia and chronic neutropenia and cancer. In addition they are also useful for treating or diagnosing various disorders associated with cell death, including myocardial infarction, stroke, neurological diseases including Alzheimer's and Parkinson's diseases, amyotrophic lateral sclerosis and spinal muscular atrophy. IFN nucleic acids and polypeptides are also useful for screening drugs and compounds which inhibit or enhance IFN activity or function and as targets for the identification of small molecules that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          08-SEP-1999;
20-OCT-1999;
20-JAN-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    determining the presence of or predisposition to a disease associated with altered levels of IFN polypeptide or polynucleotide. IFN nucleic acids, polypeptides and antibodies are useful for diagnosis, prevention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The sequence represents interferon induced nucleic acid, IFN7. IFN nucleic acids and polypeptides are useful for treating or preventing apathology associated with IFN polypeptide in a human. They are useful pathology associated with IFN polypeptide in a human.
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                                                             are immunostimulatory, immunosuppressive, or stimulate or suppress
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      or tumour cell growth in mammals,
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Sequence 3704 BP; 1168 A;

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1014 T; 0 other;

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Best Local Sim
Matches 722;
                     Homo
                                                                                                03-SEP-2001
                                                                                                                      AAH33745;
                                                                                                                                             AAH33745 standard;
                                          colorectal carcinoma;
                                                      Human; colon cancer; colon
                                                                          Human colon cancer antigen
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                                                                                                                                                                                                                                                                                           aaatgaatcaygtgtaagagagctactaaaaaggattgtgcagaaagaaaactggttctc
                                                                                                                                                                                                                                                                                                                                                                                    cctccttcagcccactctggtggacaagcttctagttagagacgtcttggataagtgcat
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                                                                                                (first entry)
                                                                                                                                             CDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19.7%;
92.9%;
                                           SS
                                                                          encoding cDNA SEQ ID NO:801.
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Pred. No. 3.3e-154;
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                                                      antigen;
                                                      diagnosis; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 3704;
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        present invention.

N.B. Pages 666 to 682 and page 7053 of the missing at time of publication, meaning no
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             inactive proteins or to supplement the patients own production of P. Additionally, N may be used to produce the colon cancer-associated Ps, by inserting the nucleic acids into a host cell and culturing the cell to express the proteins. N and P can be used in the prevention, diagnosis and treatment of colorectal carcinomas and cancers. AAH37196 to AAH37204 and AAB77789 represent sequences used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cancer antigens have cytostatic activity and can be used in gene therapy and vaccine production. N and P may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate P
                   3145
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             diagnosis and treatment of diseases associated with inappropriate P expression. For example, N and P may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of P by expressing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cancer-associated nucleic acid molecules (N) and proteins (P), where the proteins are collectively known as colon cancer antigens. The colon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 609
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29-SEP-1999;
                                                                                     3085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    useful for preventing,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        P-PSDB;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ID NO:1027 to 1052, 7921 and 7922.
               aataattcaacaaaqaaacaatacaaaaagtgqqtagaattacctatcacatttcccaat
                                                                                                                                                                                                                                                                                                                                         gccaagcattacaagaataacccatcactaataactttcctttgcaaaaactgcagtgtg
                                                                                  gtycacaaaggcttagatttyccttgtctcaaaataaggaattttgtagtggttttcaaa
                                                                                                                                                                                                      | CCAGAATTCAAGGAACTTTACATTGTAAGAGAAAACAAAGGCACTGCAAAAGGAAGTGTGCC
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                                                gtgcacaaaggcttagatttgccttgtctcaaaataaggaattttgtagtggttttcaaa
                                                                                                                   gactatcaaataaatggtgaaatcatctgcaaatgtggccaggcttggggaacaatgatg
                                                                                                                                                    gactatcaaataaatggtgaaatcatctgcaaatgtggccaggcttggggaacaatgatg
                                                                                                                                                                                     ccaqaattcaaggaactttacattgtaagagaaaacaaarcactgcaaaagaagtgtgcc
                                                                                                                                                                                                                                                      ctaqcctgttctggggaagatatccatgtaattgagaaaatgcatcacgtcaatatgacc
                                                                                                                                                                                                                                                                       gccargcattmcargaatamcccatcactaataactttcctttgcaaaaactgcagtgtg
                                                                                                                                                                                                                                                                                                                                                                                        1; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   acids encoding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              В₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2000WO-US26524.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2766; 9803pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        99US-0163280.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     9905-0157137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              229 A; 90 C; 104 G; 174 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17.2%;
98.6%;
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diagnosing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 577.2; DB 22
Pred. No. 2.6e-133;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    colon cancer-associated polypeptides,
and/or treating colorectal cancers -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rosen CA;
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acaacagcaccatctqcttgggagaaccctctcccttctctgagaaagaaagatqtcgaa 176

595

Best Loc Matches Query Match

Local

Similarity

15.1%; 99.4%;

509;

Conservative

0;

Score 507.2; Pred. No. 5.9e 0; Mismatches

.9e-116;

indels Length

Gaps

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ВП

595; 0;

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RESULT 8
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30-JUN-2000;
03-AUG-2000;
                      The present sequence is one such probe. The probes are useful for producing a microarray for predicting, measuring and displaying greynession in samples derived from human placenta. The probes are for antenatal diagnosis of human genetic disorders.
                                                                                                                                                          11dM
                                                                                                                       analyzing
                                                                                                                                                                                                                                                                                                                                                                                                                              genetic disorder;
Sequence 595
                                                                        The
                                                                                               Claim 25; SEQ ID No 9583; 654pp; English.
                                                                                                                                                                                  Penn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AA140897 standard; DNA; 595
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                                                                                                                                  Human
                                                                                                                                                                                                          (MOLE-)
                                                                                                                                                                                                                                             21-SEP-2000;
27-SEP-2000;
                                                                                                                                                                                                                                                                                                                               30-JAN-2001; 2001WO-US00663
                                                                                                                                                                                                                                                                                                                                                     09-AUG-2001.
                                                                                                                                                                                                                                                                                                                                                                              W0200157272-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                      Probe; microarray; human; placenta; antenatal diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 545
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                                                                       present invention relates to single exon nucleic acid probes (SENP).
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                                                                                                                     genome-derived single exon nucleic zing gene expression in human placen
                                                                                                                                                                                                                                                                                                                                                                                                                                                             #9583 used to measure gene expression in human placenta
                                                                                                                                                                                                         MOLECULAR DYNAMICS INC
                                                                                                                                                                                  Hanzel DK,
                                                                                                                                                                                                                              2000US-0180312.
2000US-0207456.
2000US-0608408.
2000US-0632366.
2000US-0234687.
2000US-02346359.
2000US-0236359.
BP; 141 A; 162
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                                                                                                                      in human placenta
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C; 154 G; 138
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T; 0 other;
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RESULT 9
AAI54039/c
ID AAI540
XX AAI540
XX AAI540
XX Probe
XX Probe;
KW Probe;
KW Geneti
XX Y
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30-JUN-2000;
03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
  The
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                                                    Claim 25; SEQ ID No 22725; 654pp; English.
                                                                                                            analyzing
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2000US-0207456.
2000US-0608408.
2000US-0632366.
2000US-0234687.
2000US-02346359.
2000US-0236359.
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Best Local
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                                                                                                                                                                Homo sapiens
                                                                                                                                                                                               aging process; type I interferon;
                                                                                                                                                                                                cellular senescence; terminal differentiation; growth suppression; aging process; type I interferon; cancer cell; tissue regeneration;
                                                                                                                                                                                                                                 OLD-35; OLD-64; OLD-137; OLD-139; OLD-142; OLD-175; cancerous phenotype
                                                                                                                                                                                                                                                                                                  02-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                AAA64601 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 499 BP; 116 A; 142 C; 126 G;
(UYCO ) UNIV COLUMBIA NEW YORK
                                  02-FEB-1999;
                                                                   02-FEB-2000;
                                                                                                   10-AUG-2000
                                                                                                                                  WO200046391-A2
                                                                                                                                                                                                                                                                  Nucleotide sequence of a human OLD-139 polynucleotide
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                                                                 2000WO-US02920
                                                                                                                                                                                                                                                                                                 (first entry)
                                9908-0243277
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99.3%;
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Pred. No. 4.2e-103;
0; Mismatches 3;
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are useful
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          the cancerous phenotype of a cancer cell, determining if a cell is senescent, growth arrested or terminally differentiated. They are also useful for reversing the aging process in a cell and degrading specific RNAs in a cell. The genes may also be used as a diagnostic indicator of cellular senescence, terminal differentiation and/or growth suppression and as a marker to identify drugs or small molecules that will induce or inhibit cellular senescence or terminal differentiation and type I interferons. The combination of Old-35 with other interacting proteins is useful for targeting the differentiation of specific cells. Old-35 can be used to selectively stabilize specific mkNAs containing adenoviral
                       Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3209
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   rich 3' UTRs. The OLD proteins are useful for reversing the cancerous phenotype of a cancer cell and inhibiting the growth of a cancer cell. They are also useful for regenerating tissue. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 4; Page 101; 115pp; English
breast cancer; lung cancer; cancer detection;
                                                           Novel human polynucleotide, SEQ ID NO:
                                                                                                 09-APR-2001 (first entry)
                                                                                                                                                                               AAF66485 standard; cDNA; 392
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel isolated nucleic acid encoding an OLD-35 or OLD-64 protein useful in the treatment and detection of e.g. cancer and diseases involving
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     represents an OLD-139 polynucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             181
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   specification describes OLD-35, OLD-64, OLD-137, OLD-139, OLD-142 OLD-175 proteins. The OLD nucleic acids are useful for reversing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          aattcaaggaactttacattgtaagagaaacaaagcactgcaaaagaagtgtgccgact 3028
                                                                                                                                                                                                                                                                                                            ctgacataagaatcaataaaatgattgttttactctg 3365
                                                                                                                                                                                                                                                                                                                                                                                                                                                 acaaaggcttagatttgccttgtctcaaaattagggaattttgtagtggttttcaaaaata 3148
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                                                                                                                                                                                                                                                                                                                                                  aaatactatcagttaaacatttaatatgattatgattaatgnattcattatgctncag-a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   acaaaggcllagatttgccttgtctcaaaataaggaattttgtagtggttttcaaaaata
                       cytostatic; gene therapy; colon cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             458
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97.7%;
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Pred. No. 1.4
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                      to produce probes for detection of mRNA and to produce additional copies of the polynucleotides. The probes can be used for chromosome mapping of the polynucleotide and for detection of transcription levels. Ribozymes or antisense oligonucleotides can be generated. The polynucleotides and their gene products are used as genetic or blochemical markers (e.g. in blood or tissues) that will detect the earliest changes along the carcinogenesis pathway and/or monitor the efficacy of therapies and preventive interventions. The polynucleotides, polypeptides and antibodies against them can be used in pharmaceutical compositions to treat the cancers and proliferative disorders such as neoplasia,
                                                                                                                                                2784
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         polynucleotides. The library is used to detect unitary and can genes correlated with a cancerous state of a mammalian cell and can detect colon, prostate, breast and lung cancer. The library can be used detect colon.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Library of polynucleotides for diagnosing a cancerous state mammalian cell and detecting cancer, particularly of the colprostate, comprises 3351 human polynucleotide sequences -
                                                                                                                                                                                                                                                              2664 ggttgctcacagtggttcaggagttalcgaacatgaqacagttaatgatttccqagagaa 2723
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-JUL-1999;
02-JUL-1999;
                                                                                                                                                                                             2724 gatgatgtataaagctatacattgtgttcaaaatatgaaaccagaggagtatgctcataa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence is one of 3351 sequences in a library of human polynucleotides. The library is used to detect differentially expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 9; Page 871; 1046pp; English
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Reinhard C,
                                                                                                                                                                                                                                                                                                                                                                                             Sequence
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                                                                               tgccaagcattacaagaatuacccatcactaataactttcctttgcaaaaactqcagtgt
                                                                                                                               gctagcctgttctggggaagatatccatgtaattgagaaaatgcatcacgtcaatatgac
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                                                                                                                                                                                                                                              ggttgctcacagtggttcacgagttatcgaacatgatacagttaatgatttccgagagaa
                                                                                                               gattttggaattacagatgcacagtataatggaaaagaaaatgaagaccaagagaaatat
                                               tgccaagcattacaagaataacccatcactaataactttcctttgcaaaaactgcagtgt
                                                                                                                                                                                                                                                                                                                 376;
                                                                ns LT, Escobedo J, rd C, Randazzo F, Kakov R, Drmanac S, Garcia V, Jones I
                                                                                                                                                                                                                                                                                                                                                                                             392
                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                             BP; 148
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                                                                                                                                                                                                                                                                                                                            11.0%;
97.9%;
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    Innis MA, Garcia F
Kennedy GC, Pot D,
    Dickson M, Labat I
Strache-Crain F

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                                                                                                                                                                                                                                                                                                             Score 371.2; |
Pred. No. 2.5e
0; Mismatches
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Pot D, L
Labat I,
Le-Crain B;
                                                                                                                                                                                                                                                                                                                                                                                             T; 0 other;
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D, Lamson
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                                                                 Вb
   용
                                                                                                                                Matches 367; Conserv
                                                                                                                                                                                                                                                     The present sequence represents a human expressed sequence tag (EST). The polynucleotide, which is a secreted EST, and the encoded protein are predicted to have useful biological activities which would make them suitable for treating, preventing or ameliorating medical conditions in humans and animals, although no supporting data is given. Suggested activities include nutritional activity, immune stimulating or suppressing activity, haematopoiesis regulating activity, tissue growth activity, activiny, hinhibin activity, chemotactic/chemokinetic activity, activiny, hamboritatic and thrombolytic activity, receptor/ligand activity, anti-inflammatory activity, cadherin/tumour invasion suppressor activity, tumour inhibition activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New polynucleotides encoding human secreted proteins - derived from e.g. human blood, kidney, foetal lung, placenta, testes, brain, ovary, pituitary, retina and colon cDNA libraries.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; secreted protein; expressed sequence tag; EST; haematopoiesis; tissue growth; activin; inhibin; chemotaxis; chemokinesis; haemostatic; receptor; ligand; thrombolytic; anti-inflammatory; cadherin; anti-tumour;
               1824 gaaacttotagaaataatgacaaggattcaaacttattgtcaaatgagtccaatgtcaga 1883
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; Page 277; 618pp; English.
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                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                          activity. The polynucleotide may also be useful for gene therapy
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   62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   therapy; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cgactatcacataaatggtgaaat 391
                                                              gaaacttctagaaataatgacaaggattcaaacttattgtcaaatgagtccaatgtcaga 61
3
                                                                                                                                                                                                               374 BP; 146 A; 47 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      J, Jacobs K,
Spaulding V,
                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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                                                                                                                                             10.9%;
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                                                                                                                                0;
                                                                                                                          Score 365.8; DB 20;
Pred. No. 5.4e-81;
Pred. No. 5.4e-81;
                                                                                                                                                                                                               88
                                                                                                                                                                                                               G; 93 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      мссоу лм,
                                                                                                                                                                                                             0 other;
                                                                                                                                                            DB 20;
                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Merberg D;
                                                                                                                                                            Length
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        Вþ
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                                                                          The present sequence is one of 3351 sequences in a library of human polynucleotides. The library is used to detect differentially expressed genes correlated with a cancerous state of a mammalian cell and can detect colon, prostate, breast and lung cancer. The library can be used to produce probes for detection of mRNA and to produce additional copies of the polynucleotides. The probes can be used for chromosome mapping of the polynucleotide and for detection of transcription levels. Ribozymes or antisense oligonucleotides can be generated. The polynucleotides and their gene products are used as genetic or biochemical markers (e.g. in the polynucleotides that it is the polynucleotides can be second to product the polynucleotides can be used to produce the polynucleotides can 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1944 aggaaatcgcaaagaacgtgtttgtgcagaacatttgaggaagtacaatgaggccctaca 2003
preventive interventions. The polynucleotides, polypeptides and
                            blood or tissues) that will detect the earliest changes along the carcinogenesis pathway and/or monitor the efficacy of therapies and
                                                                                                                                                                                                                                                                                                                                                            Claim 9; Page 874; 1046pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                  prostate, comprises 3351 human polynucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                         Library of polynucleotides for diagnosing a cancerous state of mammalian cell and detecting cancer, particularly of the colon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Reinhard C, I
Crkenjakov R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Williams LT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-JUL-1999;
02-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              breast cancer; lung
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel human polynucleotide, SEQ ID NO: 2255
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAF66499 standard; cDNA; 387
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2184 agatagatt 2192
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-JUN-2000; 2000WO-US18374
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cytostatic; gene therapy; colon cancer; prostate cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Garcia V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Randazzo F,
Drmanac S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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Dickson M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kennedy GC, Pot D, I
Dickson M, Labat I,
LW, Strache-Crain B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Innis MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Garcia PD, Nico,
                                                                                                                                                                                                                                                                                                                                                                                                                  sequences
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amson G, Drmanac
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Best Local Sim
Matches 364;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2182
            partic.
                      New method for generation of subtracted cDNA libraries
                                                                       Fisher
                                                                                                                                                                                04-MAY-1995
                                                                                                                                                                                                                                                                                       Melanoma differentiation associated gene mda-5 fragment
                                                                                                                                                                                                                                                                                                                 27-OCT-1995
                                                                                                                                                                                                                                                                                                                                                              AAQ90050 standard; cDNA; 301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 387 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                dysplasia and hyperplasia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antibodies against them can be used in pharmaceutical compositions treat the cancers and proliferative disorders such as neoplasia,
 partic. for identifying melanoma differentiation associated genes
for cancer study, diagnosis and therapy.
                                                WPI; 1995-178878/23.
                                                                                                                       27-OCT-1993;
                                                                                                                                  30-SEP-1994;
                                                                                                                                                         24-OCT-1994;
                                                                                                                                                                                                        W09511986-A.
                                                                                                                                                                                                                             Homo sapiens
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                                                                                                                                                                                                                                                     melanoma
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                                                                                              AIND (
                                                                        PΒ,
                                                                       Jiang H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                               COLUMBIA NEW YORK
                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                      cell;
                                                                                                                     94US-0316537.
93US-0143576.
                                                                                                                                                         94WO-US12160
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96.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                        386
and therapy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 356.8; DB 2
Pred. No. 9.4e-79;
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Best Local Similarity
Matches 297; Conserv
                                                   04-FEB-2000;
26-MAY-2000;
30-JUN-2000;
03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Clinical Research Centre, Watford Road, Harrow, Middx., HAI, England. It represents a novel IFN-gamma inducible gene which also displays increased expression during terminal cell diffn. in HO-1 cells. It represents a cytokine-, and terminal differentiation-inducible gene displaying increased expression in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1652
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mda-5 cDNA is novel. It has sequence homology to a Homo sapiens putatively transcribed partial sequence; Accession no. AAZ20545 the UK-HGMP, MRC Human Genome Mapping Project Resource, Centre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1712
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1592 acaaaccagtgattccccttcctcagatactgggactaacagcttcacctggtgttgyag
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             all melanomas, in select carcinomas, in normal skin fibroblasts in both normal cerebellum and \ensuremath{\mathsf{GBM}}.
                                                                                                                                                                                   09-AUG-2001
                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                     cervical cancer; ss
                                                                                                                                                                                                                                                                                  Probe; human; microarray; gene expression; cervical epithelial cell;
                                                                                                                                                                                                                                                                                                             Probe #9846 for gene expression analysis in human cervical cell sample
                                                                                                                                                                                                                                                                                                                                           12-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                   AA119913 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1832 tagaaataatgacaaggaltcaaacttattgtcaaatgagtccaatgtcagattttgga 1890
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 46; Page 216; 421pp; English
           (MOLE-) MOLECULAR DYNAMICS INC
                                                                                                                                                       30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       240
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PI Penn SG, Hanzel DK, Chen W, Rank DR;

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Human genome-derived single exon nucleic acid probes useful for pr analyzing gene expression in human cervical epithelial cells.

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Claim 25; SEQ ID No 9846; 487pp; English.

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The present invention relates to human single exon nucleic acid probes (C (SENP). The present sequence is one such probe. The SENPs are derived (C (SENP). The present sequence is one such probe. The SENPs are derived (C (SENP). The present sequence is one such probe. The serviced in a single exon the probe of the can be used for measuring human gene expression in a cc variety of the can be used for measuring human gene expression in a cc variety of the can be used in a grading and/or staging (C (SENP)). The probes are therefore useful in grading and/or staging (C (SENP)) and the cervix, notably cervical cancer.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO (at ftp. wipo.int/pub/published_pct_sequences.)

Sequence 292 BP; 108 A; 31 C; 71 G; 82 T; 0 other:
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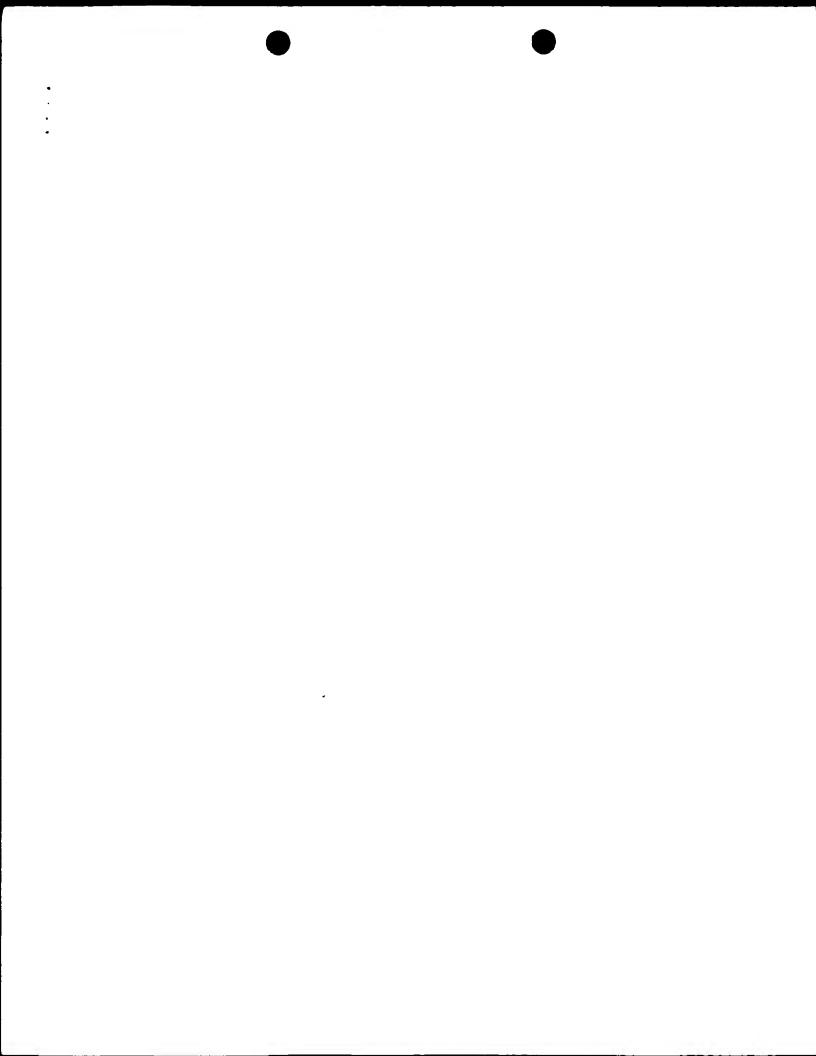
Query Match 8.4%; Score 281.4; DB 22; Length 292; Best Local Similarity 97.9%; Pred. No. 4.1e-60;

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ORGANISM REFERENCE AUTHORS TITLE RESULT AF095844 LOCUS JOURNAL REFERENCE ACCESSION DEFINITION AUTHORS TITLE Unpublished
2 (bases 1 to 3380)
Kang,D.-C. and Fisher,P.B.
Direct Submission Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 3380)
Kang,D.-C. and Fisher,P.B.
Kang,D.-C. and Fisher,P.B. AF095844 3380 bp mrNA PRI 26-NOV-2000 Homo sapiens melanoma differentiation associated protein-5 (MDA5) mRNA, complete cds. AF095844 Homo sapiens AF095844.1 G1:11344593 inducible gene of limited homology to RNA helicase human

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2 (bases 1 to 3373)
Cocude,C., Kolesnitchenko,V.,
Capron,A. and Bahr,G.M.
                                                               Mammaila; Eutheria; Primates;
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Cocude.C
                                                    Cocude, C., Kolesnitchenko, V., Capron, A. and Bahr, G.M.
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Submitted (09-JAN-2001) Laboratoire d'Immunologie Moleculaire
l'Infection et de l'Inflammation, Institut Pasteur de Lille, 1
du Professeur Calmette BP 245, Lille 59019, France
Location/Qualifiers
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         NIH-MGC Project URL: http://mgc.nci.nih.qov
Contact: MGC help desk
                                                       Direct Submission
Submitted (11-MAY-2001)
                                                                                                                                                        Homo sapiens,
BC007966
Email: cgapbs-r@mail.nih
                                               Gene Collection (MGC),
                                                                              Strausberg,R
                                                                                                Mammalia;
                                                                                                         Eukaryota;
                                                                                                                    Homo sapiens
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                                       Institute,
                                                                                       (bases 1 to 1213)
                                                                                              Eutheria;
                                       31 Center
                                                                                                                                               GI:14044090
                                                                                                                                                                 1213 bp
, clone I
                                                                                               Chordata;
Primates;
                                                                                                                                                                   IMAGE: 4304805,
                                       Drive,
                                                Cancer
                                               National Institutes of Health, Mammalian Nancer Genomics Office, National Cancer
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                                       Room
                                                                                                Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                       11A03,
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                                      Bethesda, MD
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Gaithersburg, Maryland;
Web site: http://www.
Contact: nisc_mgc@nh
Shevchenko,Y., Wetherby,K.E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
series: IRAL Plate: 20 Row: p Column: 10
This clone was selected for full length sequencing because it
passed the following selection criteria: Hexamer frequency ORF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: nlsc_mgc@nhgrl.nih.gov
Shevchenko,Y., Wetherby,K.D., Beckstrom-Sternberg,S.M.,
Benjamin,B., Blakesley,R.W., Bouffard,G.G., Brinkley,C., Brooks,S.,
Benjamin,B., Guan,X., Gupta,J., Ho,S.-L., Karlins,E., Legaspi,R.,
Lim,M., Maduro,Q.L., Maslello,C., Mastrian,S.D., McCloskey,J.C.,
McDowell,J., Pearson,R., Snyder,B., Stantripop,S., Thomas,P.J.,
Tiongson,E.E., Touchman,J.W., Tsurgeon,C., Vogt,J.L., Walker,M.A.,
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cDNA Library Preparation: Rub
cDNA Library Arrayed by: The
DNA Sequencing by: National I
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/tissue_type="Uterus, leiomyosarcoma"
/clone_lib="NHH_MGC 46"
/clone_lib="NHH_MGC 46"
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247 c 285 g
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/db_xref="taxon:9606"
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Pred. No. 3e-2:
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The I.M.A.G.E. Consortium (LLNL)
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Contact: MGC help desk
Email: cgapbs-r@mail.nih.gov
                                                                              Submitted (28-PEB-2001) National Institutes of Health, Mammalian Gene Collection (MGC), Cancer Genomics Office, National Cancer Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 2050)
                                                                                                                              Direct Submission
                                                                                                                                               Strausberg,R.
                                                                                                                                                                                                             Mus musculus
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                                                                                                                                                                                                                                                                                           Mus musculus,
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Tissue Procurement: Gilbert Smith,
                                                                                                                                                                                                                                                                                                            2050 bp
                                                                                                                                                                                                                                                                                         clone IMAGE: 3495361, mRNA, partial cds
                                                                                                                                                                                                                                                                                                          mRNA
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FEATURES
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CTACAAATCAACGACACGATCCGAATGATTGATGCGTATAGCCACCTGGAGACATTCTAC
                                                                                                          aaaaaaggaaatcgcaaagaacgtgtttgtgcagaacatttgaggaagtacaatgaggcc
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                                                                                                                                                                                                                                                                                                                                                                         caaatacaggagccatgcaagaagtttgccattgccagatgcaaccagagaagatccattt 1818
                        TCAGATTTTGGAACCCAACATTATGAGCAGTGGGCCATTCAAATGGAGAAAAAAGCTGCT
                                                                                                                                                                                                                AAAGAGAAACTACTAGAAATTATGGCAAGCATTCAGACTTACTGCCAAAAAAAGTCCAATG
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                                                                                      AAAGACGGAAATCGCAAAGATCGCGTCTGTGCAGAACATTTGAGGAAGTACAACGAAGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Clone distribution: MGC clone distribution information can be fo through the I.M.A.G.E. Consortium/LLML at: http://lmage.llnl.gov Series: IRAK Plate: 8 Row: k Column: 18 This clone was selected for full length sequencing because it passed the following selection criteria: Hexamer frequency ORF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: (Di
Dickson, M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Sequencing Group at the Stanford Human Genome
Center, Stanford University School of Medicine, Stanford, CA 94:
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VNDFREKMMYKAINRVQNMKPEEYAHKILELQVQSILEKKMKVKRSIAKQYNDNPSLI
TLLCKNCSMLVCSGENIHVIEKMHHVNMTPEFKGLYIVRENKALQKKFADYQTNGEII
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/clone_lib="NCI_CGAP_Lu29"
/lab_host="DH10B"
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STYALSQWIMENAKFAEYGYKAHHLIGAGHSSEYKPMTQTEQKEYISKFRTGEINLLI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LEIMAS I QTYCQKSPMSDFGTQHYEQWA I QMEKKAAKDGNRKDRYCAEHLRKYNEALQ
INDTIRMIDAYSHLETFYTDEKEKKFAVLNDSDESDDEASSCNDQLKGDYKKSLKLDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /codon_start=1
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/protein_id="AAH04031.1"
/db_xref="01:13278456"
/translation="ANLDAFTIKTVKENLGQLKHQIKEPCKKFVIADDTRENPFKEKL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /tissue_type="Mammary tumor metastatized to lung. Tumor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /db_xref="taxon:10090"
/clone="IMAGE:3495361"
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(Dickson, Mark) mcd@paxil.stanford.edu
M., Schmutz, J., Grimwood, J., Rodriquez, A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note="Vector: pCMV-SPORT6"
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                                                                                                        TTTGCTGATTATCAGACCAATGGAGAGATTATCTGCAAGTGTGGCCAGGCTTGGGGAACA
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Mammalla; Eutherla; Primates;
1 (bases 1 to 1258)
Peyman,J.A., da Silva,A., Hoch
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                                                             /protein_id="CAC35442.1"
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KRTVQKENWFSAFI.NVLRQTGNNELVQELIFGSDCSESNAETENLSQVDGPQVEEQLLS
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Peyman, J.A., da Silva, A., Hochman, P. and Hsu, A. Interferon induced polynucleotides and proteins Patent: WO 0118208-A 9 15-MAR-2001; Curagen Corporation (US): BIOGEN, INC. (US)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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a 709 c 806 g 1011 t
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                           Score 663; DB 6;
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KRIVQKENWFSAFLNVLRQTGNNELVQELTGSDCSESNAGICNFTEEDSSNSA"
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SGNWQAVELLLSTLEKGVWHLGWTREFVEALRRTGSPLAARYMNPELTDLPSPSFENA
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                                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Homo sapiens"
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        Direct Submission
Submitted (30-SEP-2000) Department of Genetics, Washington
University, 4444 Forest Park Avenue, St. Louis, Missouri 6
On Mar 15, 2000 this sequence version replaced gi:5923734
                                                                                                                       Direct Submission
Submitted (15-MAR-2000)
University School of Med
                                                                            Waterston, R.
                                                                                                                                                                                                        MO 63108,
                                                                                                                                                                                                                   Submitted (25-SEP-1999) Genome Sequencing Center, Washington University School of Medicine, 4444 Forest Park Parkway, St.
                                                                                                                                                                                                                                                    Direct Submission
                                                                                                                                                                                                                                                                                                Unpublished
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Center: Washington University Genome

Sequencing Center

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FEATURES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NOTICE: This sequence may not represent the entire insert of this clone. It may be shorter because we only sequence overlapping clone sections once, or longer because we provide a small overlap between neighboring data submissions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The RPCI-11 human BAC library was made from the blood of one male donor, as described by Osoeqawa, K., Woon, P.Y., Zhao, B., Frengen, E., Tateno, M., Catanese, J.J. and de Jong, P.J. (1998) An improved approach for construction of bacterial artificial chromosome libraries. Genomics 51:1-8. The clone may be obtained either from proceedings of the control of the clone may be obtained either from the processor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mapping information for this clone was provided by Dr. John D. McPherson, Department of Genetics, Washington University, St. Louis MO. For additional information about the map position of this
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The clone sequenced to the left is RP11-576116, 200 bp overlap; the clone sequenced to the right is RP11-204024. Actual start of this clone is at base position 126545 of RP11-576116; actual end is at base position 81443 of RP3-404K21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MAPPING INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The sequence RP11-214A4 from base postion 23333 to 23415 was derived from per off clone DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Research Genetics, Inc. (http://www.resgen.com) or Pieter de Jong
and coworkers at the Roswell Park Cancer Institute
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOURCE INFORMATION:
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Contact: sapiens@watson.wustl.edu
Summary Statistics
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695. .726
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2364. .2505
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763. .950
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/clone_lib="RPCI-11"
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8919, .8968
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7585. .7694
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11237. .11409
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repeat_region

/rpt_family="MaLR" 27897. .28473

/rpt_family="Mal.R"

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29139. .29270
/note-"similar to Mus musculus EST W29262 (NID:g1309410)
mc22c08.r1"
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Mammalia; Eutheria; F
1 (bases 1 to 392)
Williams,L.T., Escobe
                           Eukaryota; Metazoa: Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 387) Williams, L.T., Escobedo, J., Innis, M.A., Garcia, P.D., Klinger, J., Kassam, A., Reinhard, C., Randazzo, F., Kennedy, G.C., Pot, D., Lamson, G., Drmanac, R., Crkenjakov, R., Drmanac, S., Dickson, M., Labat, J., Leshkowitiz, D., Kita, D., Garcia, V. and Strache-Crain, B. Human genes and gene expression products
Patent: WO 010258-A 2255 11-JAN-2001;
                                                                                                                                                                                                                                                                                Sequence 2255
AX071783
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Lamson,G., Drmanac,R., Crkenjakov,R., Drmanac,S., Dickson,M.,
Labat,I., Leshkowitiz,D., Kita,D., Garcia,V. and Strache-Crain,B
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Pred. No. 3.le-68;
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                                                                                                                                                               Direct Submission
Direct Submission
Submitted (05-JUN-1999) Genome Sequencing Center, Washington Submitted (05-JUN-1999) Genome Sequencing Center, Washington Submitted (05-JUN-1999) Genome Sequencing Center, Washington
5 MO
                                        Submitted (22-OCT-1999) Genome Sequencing Center, Washington University School of Medicine, 4444 Forest Park Parkway, St. Louis,
                                                                                                                                                                                                                                                                                                                    Cotton, M., Maupin, R., Hawkins, M. and Harkins.
The sequence of Homo sapiens BAC clone RPll:
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Sulston, J.E. and Waterston, R.
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Homo sapiens BAC clone
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                                                                                                               Waterston, R.H.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The RPCI-II human BAC library was made from the blood of one male donor, as described by Osoegawa, K., Woon, P.Y., Zhao, B., Frengen, E., Tateno, M., Catanese, J.J., and de Jong, P.J. (1998) An improved approach for construction of bacterial artificial chromosome libraries. Genomics 51:1-8. The clone may be obtained either from Research Genetics, Inc. (http://www.resgen.com) or Pieter de Jong and coworkers at the Roswell Park Cancer Institute
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (02-OCT-2000) Department of Genetics, Washington University, 4444 Forest Park Avenue, St. Louis, Missouri 6 On Oct 22, 1999 this sequence version replaced gi:5103889.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The clone sequenced to the left is RP11-178A14. Actual start of this clone is at base position 1 of RP11-576I16; actual end is at base position 1636B1 of RP11-576I16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mapping information for this clone was provided by Dr. John D. McPherson, Department of Genetics, Washington University, St. 1 MO. For additional information about the map position of this
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NOTICE: This sequence may not represent the entire insert of this clone. It may be shorter because we only sequence overlapping clone sections once, or longer because we provide a small overlap
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Waterston, R.
Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MAPPING INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This sequence was finished as follows unless otherwise noted:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    between neighboring data submissions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NEIGHBORING SEQUENCE INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOURCE INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sequence, see http://genome.wustl.edu/gsc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    restriction digest.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (http://bacpac.med.buffalo.edu)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Web site: http://genome.wustl.edu/gsc
Contact: sapiens@watson.wustl.edu
Summary Statistics
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                  /rpt_family="(CA)n"
5417. .5648
                                                                /rpt_family="AT_rich" 5263, .5299
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1536. .1694
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1271. .1298
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216 .322
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                                                                                                                                                                             /rpt_family="Alu"
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/chromosome="2"
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9386. .9586
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8702. .9187
                                                /note="match to EST AI589439 (NID:g4598487) tr76a08.x1"
21204, .21219
                                                                            /rpt_family-"(T)n"
21204. .21219
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20851. .20910
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20563. .20850
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16625.
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21204. .21219
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15713, .15993
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14742. .14887
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13833. .14130
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13109. .13133
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11136.
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10580. .10739
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6. .15695
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5. .16997
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4. .13419
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VERSION
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                                                                                                                                                           Unknown
                                                                      Patent: US 5643761-A 7 01-JUL-1997;
                                                                                  Method for generating a subtracted cDNA library and uses generated library
                                                                                                              1 (bases 1 to 301)
Fisher, P.B. and Jiang, H.
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23343. .23397
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             /organism="unknown"
61 c 56 g
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24730. .25234
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19. .23983
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Best Local Similarity 99.3%;
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                                             AACTGTTAAAGAAAACCTTGATCAACTGAAAAACCAAATACAGGAGCCATGCAAGAAGTT
                                                                       aactgttaaaqaaaaccttgatcaactgaaaaaccaaatacaggagccatgcaagaagtt 1784
                                                                                                     AGCCAAAGCTGAAGAACACATTTTAAAACTATGTGCCAATCTTGATGCATTTACTATTAA 182
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Sibson,D.R. and Gross,J.
HUMAN NUCLEIC ACID FRAGMENTS, ISOLATED FROM BRAIN ADRENAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 (bases 1 to 377)
Sibson,D.R. and Hadfield,K.M.
HUMAN NUCLEIC ACID FRAGMENTS ISOLATED FROM BRAIN, ADRENAL TISSUE,
PLACENTA OR BONE NARROW AND THEIR USE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent: EP 0587279-A 240 16-MAR-1994; MEDICAL RES COUNCIL (GB)
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length DB		ID 01	Description
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2	687	20.4	729	11	BF983236	
ω	652.8	19.4	827	11	BG741146	-
4	651.8	19.4	781	11	BF686405	
_U	622.8	18.5	1041	11	BE882040	
6	618.4	18.4	1197	12	AK018602	
c 7	531.4	15.8	536	10	A1718277	AI718277 as51e06.x
o 8	520.4	15.5	537	10	A1806204	AI806204 wf26c12.x
c 9	515.4	15.3	521	10	AI922705	
c 10	511.2	15.2	552	10	AA134958	AA134958 zo23f04.s
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Z20545 HSAAACJNK S AA177334 mt17g10.r BF124647 601761004	uf51f08. ud83b06. uf31e11.	0	114a01.r1 BB284625 RPCI11-7	AI906351 RC-BT108- AI783469 tz35d07.x		o		RC6-		- 00	BG967530 602833306 AW580584 CM3-HT048	BG391418 602417507 AW779876 hn89c07.x		9567	AW152541 xf76b02.x

ALIGNMENTS

ce	FEATURES		REFERENCE AUTHORS TITLE JOURNAL COMMENT	VERSION KEYWORDS SOURCE ORGANISM	RESULT 1 BF337464 LOCUS DEFINITION ACCESSION
/organism."Homo sapiens" /organism."Homo sapiens" /db_xref="taxon:9606" /clone='IMAGE:4183126" /clone='Ib="MCI_CGAP_Brn64" /tissue_type="glioblastoma with EGPR amplification"	nctp://lmage.lini.gov Plate: LLAM9498 row: n column: 23 High quality sequence start: 5 High quality sequence stop: 695. Location/Qualifiers 1 1038	Email: cgapbs-remail.nih.gov Tissue Procurement: David N. Louis, M.D. cDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:	l (bases i to 1035) NIH-MGC http://mgc.nci.nih.gov/. National Institutes of Health, Mammalian Gene Collection (MGC) Onpublished (1999) Contact: Robert Strausberg, Ph.D.	BF337464.1 GI:11283715 EST. human. Homo Sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi; Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.	BF337464 1035 bp mRNA EST 22-NOV-2000 602035195F1 NCI_CGAP_Brn64 Homo sapiens cDNA cione 1MAGE:4183126 5', mRNA sequence. BF337464

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                                                        BF983236 729 bp
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/note="Organ: brain; Vector: pCMV-SPORT6; Site_1:
SIte_2: Sall; Cloned unidirectionalPORT6 Primer: 01
Average insert size 1.57 kb. Constructed by Life
Technologies. Note: this is a NCI_CGAP Library."
a 221 c 243 g 218 t
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IMAGE: 4397083
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                                                                  atgggcagtgattcaggcaccatgggaagtgattcagatgaagagaatgtggcagcaaga
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Plate: LLAM10097 row: a column:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Robert Strausberg, Ph.D.
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1 (bases 1 to 729)
NIH-MGC http://mgc.nci.nih.qov/
National Institutes of Health, |
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/lab_host="DH10B (phage resistant)"
/note="Organ: small intestine; Vector: pCMV-SPORT6;
Site_1: Not1; Site_2: Sal1; Cloned unidirectionally;
oligo-dT primed. Average insert size 1.767 kb. Library
enriched for full-length clones and constructed by Life
Technologies, Note: this is a NIH_MGC Library."
a 137 c 184 g 168 t
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/clone="IMAGE:4397083"
/clone_lib~"NIH_MGC_88"
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Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Pred. No. 7.8e-131;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mRNA sequence.
BG741146
BG741146.1 GI
                                                                                                                                                                                                                                                                                                                                                                                                                 Email: cgapbs-r@mail.nih.gov
Tissue Procurement: James Cleaver, M.D.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) D
Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
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602631817F1 NCI_CGAP_Skn3 Homo sapiens cDNA clone
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NIH MGC http://mgc.nci.nih.gov/.
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Plate: LLAM10630 row: b column:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           National Institutes of Health, Mammalian
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                                                                                                                                                                                                 254
                                                                                                                                                                                                                                                                                                                                                                             quality sequence stop:
                                                                                                          Conservative
                                                                                                                                                                                        /organism="Homo sapiens"
/ob_xref="taxon:9606"
/clone="MAGE:4776869"
/clone="MAGE:4776869"
/clone="Del'NOTECGAP_Skn3"
/lab_host="DH10B (T1 phage-resistant)"
/note="organ: skin; Vector: pCMV-SPORT6; Site_1: NotI;
Site_2: Sall; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.5kb. Library constructed by Life
Technologies. Note: this is a NCI_CGAP Library."
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                                                                                                                        19.4%;
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                                                                                                         Score 652.8; DB 11;
Pred. No. 8.2e-124;
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                                                                                                   Eukaryoia; Metazoa; Chordata; Craniata; Mammalia; Eutheria; Primates; Catarrhini l (bases 1 to 781)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
                                              Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
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602143786F1 NIH_MGC_46
                                                                          Contact: Robert Strausberg, Ph.D
                                                                                           Unpublished (1999)
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CDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium () DNA Sequencing by: Incyte Genomics, Inc.
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                                                                                                                                                                                       Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                             clone IMAGE: 4304805
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                      agtcttggacataacag-caacatgggcagtgattcaggcaccatgggaagtgattcaga
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                                                                                          Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
plate: LLCM170 row: 1 column: 22
High quality sequence stop: 731.
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/lab_host="DH10B (phage-resistant)"
/note="Organ: uterus; Vector: poTB7; Site_1: XhoI; Site_2:
EcoRI; cDNA made by oligo-dT priming. Directionally cloned
into EcoRI/XhoI sites using the following 5' adaptor:
GCACGAC(G). Size selected >500bp for average insert size
1.8kb. Library constructed by Ling Hong in the laboratory
of Gerald M. Rubin (University of California, Berkeley)
using ZAP-cDNA synthesis kit (Stratagene) and Superscript
ILRT (Life Technologies). Note: this is a NIH_MGC
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/clone_11b="NIH_MGC_46"
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/db_xref="taxon:9606"
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2222 aaatgttgaaaaggetggetgaaaaeeeagaatatgaaaatgaaaagetgaeeaaattaa
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                                                                                                                                                                                                                                                                                                                                                                     121
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                                                                                                        aacctttgaaactggatgaaacagatagatttctcatgactttatttttttgaaaacaata 2221
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Tissue Procurement: ATCC
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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/db_xref="taxon:9606"
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/tissue_type="leiomyosarcoma"
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Shibata,K., Itoh,M., Alzawa,K., Nagaoka,S., Sasaki,N., Carninci,P., Konno,H., Akiyama,J., Nishi,K., Kitsunai,T., Tashiro,H., Itoh,M., Suni,N., Ishii,Y., Nakamura,S., Hazama,M., Nishine,T., Harada,A., Yamamoto,R., Matsumoto,H., Sakaguchi,S., Ikegami,T., Kashiwagi,K., Fujiwake,S., Inoue,K., Togawa,Y., Izawa,M., Ohara,E., Watahiki,M., Yoneda,Y., Ishikawa,T., Ozawa,K., Tanaka,T., Matsuura,S., Kawai,J., Okazaki,Y., Muramatsu,M., Inoue,Y., Kira,A. and Hayashizaki,Y., Okazaki,Y., Muramatsu,M., Inoue,Y., Kira,A. and Hayashizaki,Y., RIKEN integrated sequence analysis (RISA) system--384-format
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Carninci, P., Shibata, Y., Hayatsu, N., Sugahara, Y., Shibata, K., Itoh, M., Konno, H., Okazaki, Y., Muramatsu, M. and Hayashizaki, Y.
                                                                                                                                                                                                                                                                                                              Normalization and subtraction of cap-trapper-selected cDNAs to prepare full-length cDNA libraries for rapid discovery of new genes Genome research. 10 (10), 1617-1630 (2000)
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Mus musculus adult male cecum cDNA, RIKEN full-length
library, clone:9130009C22, full insert sequence.
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High-efficiency full-length cDNA cloning
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Arakawa,T., Carninci,P., Fukuda,S., Fukunishi,Y., Furuno,M., Hanagaki,T., Hara,A., Hayatsu,N., Hiramoto,K., Hiraoka,T., Hori,F., Imotani,K., Ishii,Y., Itoh,M., Izawa,M., Kato,H., Kawai,J., Kojima,Y., Konno,H., Kouda,M., Koya,S., Kurihara,C., Matsuyama,T.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  prepare mouse tissues. First strand cDNA was primed with a primer [5' GAGAGAGAGGATCCAAGAGCTCTTTTTTTTTTTTTTTVN 3'], cDNA was prepared by using trahalose thermo-activated reverse transcriptase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cDNA library was prepared and sequenced in Mouse Genome Encyclopedia Project of Genome Exploration Research Group in Rike Genomic Sciences Center and Genome Science Laboratory in RIKEN.

Division of Experimental Animal Research in Riken contributed to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          URL:http://genome.gsc.riken.go.jp/, Tel:81-45-503-9222;
Fax:81-45-503-9216)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Miyazaki,A., Nishi,K., Nomura,K., Numazaki,R., Ohno,M., Okazaki,Y., Okido,T., Owa,C., Saito,H., Saito,R., Sakai,C., Sakai,K., Sano,H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Please visit our web site (http://genome.gsc.riken.go.jp/) for further details.
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                                  /note="putative"
214 c 256
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1197
                                                                                                                                                                             GLDLPCLKIRNFVVNFKNNSPKKQYKKWVELPIRFPDLDYSEYCLYSDED*
                                                                                                                                                                                                    RVQNMKPEEYAHKILELQVQSILEKKMKVERSIAKQYNDNPSLITLLCKNCSMLVCSG
ENIHVIEKMHHVNMTPEFKGLYIVRENKALQKKFADYQTNGEIICKCGQAWGTMNVHK
                                                                                                                                                                                                                                                        /codon_start=1
/protein_id="BAB31303.1"
/db_xref="G1:12858394"
/translation="MVQARGRARADESTYVLVTSSGSGVTEREIVNDFREKMMYKAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                /tissue_type="cecum"
/clone_lib="RIKEN_full-length enriched mouse cDNA library"
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/db_xref="MGD:MGI:1904143"
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/strain="C57BL/6J"
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105. .737
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Best Local Similarity 83.7%;
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                                                                                                                                                         TATGATTTAATTGTTGTTTTCGCTACACTACTGAGCTAACTCTAGAATATGTAGA
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IMAGE:2320738 3', mRNA sequence.
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Hillier,L., Allen,M., Bowies,L., Dubuque,T., Geisel,G., Krizman,D., Kucaba,T., Lacy,M., Le,N., Lennon,G., Marra, J., Moore,B., Schellenberg,K., Steptoe,M., Tan,F., Thei White,Y., Wylie,T., Waterston,R. and Wilson,R. WashU-NCI human EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Seq primer: -40UP from Gibco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (1997)
Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                    High quality sequence stop: 481.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Washington University School of Medicine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: est@watson.wustl.edu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Forest Park Parkway, Box
Barstead."
a 93 c
                                                                                                                                               3']; double-stranded cDNA was ligated to Eco RI adaptors [5' AATTCGGATCGAAC 3' and 5' GTTGGATCGG 3'], digested with Not I and cloned into the Not I and Eco RI sites of
                                                                                                                                                                                                                                                                                                                          /db_xref="taxon:9606"
/clone="IMAGE:2320738"
/clone_lib-"Barstead aorta HPLRB6"
                                                 the modified pT7T3 vector. Library constructed by Bob
                                                                                                                                                                                                                                                                                                   /sex-"male"
                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Homo sapiens"
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2833 aagagaaatattgccaagcattacaagaataacccatcactaattaactttcctttqcaaa 2892 acatttcccaatcttgactattcagaatgctqtttatttagtgatgaggattagcacttg gtcaatatgaccccagaattcaaggaactttacattgtaagagagaaacaaagcactgcaa aactgcagtgtqctagcctgttctggggaaqatatccatqtaattgagaaaatqcatcac 2952 GTCAATATGACCCCAGAATTCAAGGAACTTTACATTGTAAGAGAAAAACAAAACACTGCAA AAGAGAAATATTGCCAAGCATTACAAGAATAACCCATCACTAATAACTTTCCTTTGCAAA 477 $\tt GTGGTTTTCAAAAATAATTCAACAAAGAAACAATACAAAAGTGGGTAGAATTACCTATC$ GGAACAATGATGGTGCACAAAGGCTTAGATTTGCCTTGTCTCAAAAATAAGGAATTTTGTA AAGAAGTGTGCCGACTATCAAATAAATGGTGAAATCATCTGCCAAATGTGGCCAGGCTTGG Conservative 15.8**%;** 99.8**%**; 0; Mismatches Score 531.4; DB 1 Pred. No. 6.6e-99; DB 10; Length 536; Indels 0; Gaps 3072 3012 297 417 0,

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3012 aaagaagtgtgccgactatcaaataaatggtgaaatcatctgcaaatgtggccaggcttg 3071
                                                                                                                        116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTGAAGATTCTTTTAAAATACTATCAGTTAAACATTTAATATGATTATGATTAATGTAT
                                           | CGTCAATATGACCCCAGAATTCAAGGAACTTTACATTGTAAGAGAAAACAAAACACTGCA 359
                                                                                  cgtcaatatgaccccagaattcaaggaactttacattgtaagagaaaacaaagcactgca 3011
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tumor Gene 1000.
Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AI806204 537 bp mRNA EST 19-D wf26C12.x1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:2356726 3', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 537)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap. National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EST
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AI806204.1 GI:5392770
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                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                   /note-"Organ: pooled; Vector: pT773D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; Equal amounts of plasmid DNA from three normalized libraries (fetal lung NbHL19W, testis NHT, and B-cell NCI_CGAP_GCB1) were mixed, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from pools of 5,000 clones made from the same 3 libraries. The pools consisted of I.M.A.G.E. clones 297480-302087, 682632-687239, 726408-728711, and 729096-731399. Subtraction by Bento Soares and M. Fatima Bonaldo. "
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /clone_lib="Soares_NFL_T_GBC_S1"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /clone="IMAGE:2356726"
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                                                                                                                                                                                                                                                                                                                  15.5%;
                                                                                                                                                                                                                                                                                             Score 520.4; DB 1
Pred. No. 1.2e-96;
0; Mismatches 1
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2849 agcattacaagaataacccatcactaataactttcctttgcaaaaactgcagtgtgctag 2908

521 AGCATTACAAGAATAACCCATCACTAATAACTTTCCTTTGCAAAAACTGCAGTGTGCTAG 462

Query Match Best Local

516;

Conservative

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Similarity

15.3%; 99.8%;

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                                                                                                                                                                                                                                                                                  www-bio.llnl.gov/bbrp/image/image.html
Seg primer: -400P from Gibco
High quality sequence stop: 410.
                                                                                                                                                                                                                                                                                                                                                DNA Sequencing by: Washington University Genome Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota: Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                             Life Technologies catalog #: 11548-013
                                                                                                                                                                                                                                                                                                                                                                                                                                Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tumor Gene Index
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   National Cancer Institute, Cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
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              /note="Organ: pancreas: Vector: pCMV-SPORT6; Site_1: SalI;
Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.72 kb. Life Technologies catalog #:
11548-013"
a 90 c 82 g 195 t
                                                                                                                  /db_xref="taxon:9606"
/clone="IMAGE:2455047"
/clone_lib="NCI_CGAP_Pan1"
/tissue_type="adenocarcinoma"
/lab_host="DH10B"
                                                                                                                                                                                                                        /organism="Homo sapiens"
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                                                                                                                                                                                                                                                                                                                                                                                 Fax: 314 286 1810
Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. 1 Tel: 314 286 1800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Underwood,K., Wohldmann,P., Waterston,R., Wilson,R. and Marra,M. Generation and analysis of 280,000 human expressed sequence tags Genome Res. 6 (9), 807-828 (1996)
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/organism."Homo sapiens"
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/db_xref.='taxon:9606"
/clone_"IMAGE:587743"
/clone_lib-"Stratagene_co
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            Dias Neto, E., García Correa, R., Verjovski-Almeida, S., Briones, M.R., Nagai, M.A., da Silva, W. Jr., 2ago, M.A., Bordin, S., Costa, F.F., Goldman, G.H., Carvalho, A.F., Matsukuma, A., Baia, G.S., Simpson, D.H., Brunstein, A., deoliveira, P.S., Bucher, P., Jongeneel, C.V., O'Hare, M.J., Soares, F., Brentani, R.R., Reis, L.F., de Souza, S.J. and
                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 546)
Simpson, A.J.
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/cell_line="TB4 carcinoma cell line"
/lab_host="SOLR cells (Kanamycin resistant)"
/note="Organ: colon; Vector: pBluescript SK-; Site_l:
/note="Organ: colon; Cloned unidirectionally. Primer:
ECORI; Site_2: XhoI; Cloned unidirectionally. Primer:
Oligo dT. T-84 colonic epithelial cell line. Average
insert size: 1.0 kb; Uni-ZAP XR Vector; -5' adaptor
sequence: 5' GAATTCGGCACGAG 3' -3' adaptor sequence: 5'
CTCGAGTTTTTTTTTTTTTTTTTTTTTTTTTTTTTT"
a 97 c 90 g 206 t 2 others
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TITLE

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                                                               ttcccaatcttgactattcagaatgctgtttatttagtgatgaggattagcacttgattg 3256
                                                                                                                                 ttttcaaaaataattcaacaaagaaacaatacaaaaagtgggtagaattacctatcacat 3196
                                                                                                                                                                                CAATGATGGTGCACAAAGGCTTAGATTTGCCTTGTCTCAAAATAAGGAATTTTGTAGTGG
                                                                                                                                                                                                 caatgatggtgcacaaaggcttagatttgccttgtctcaaaataaggaattttgtagtgg 3136
                                                                                                                                                                                                                                                ATATGACCCCAGAATTCAAGGAACTTTACATTGTAAGAGAAAACAAAGCACTGCAAAAGA
                                                                                                                                                                                                                                                                                                                                 atatgaccccagaattcaaggaactttacattgtaagagaaaacaaagcactgcaaaaga 3016
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TCATCAGATTTTGGAATTACAGATGACAAGTATAATGGAACAAGAAAATGAAAACCAAGA 70
                                              TTCCCAATCTTGACTATTCAGAATGCTGTTTATTTAGTGATGAGGATTAGCACTTGATTG
                                                                                                                TTTTCAAAAATAATTCAACAAAGAAACAATACAAAAAGTGGGTAGAATTACCTATCACAT 429
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      530;
This sequence was derived from the FAPESP/LICR Human Cancer Genome Project. This entry can be seen in the following URL (http://www.ludwig.org.br/scripts/gethtml2.pl?tl=&t2=RC3-UT0064-170 800-021-e06&t3=2000-08-17&t4-1) seq primer: puc 18 forward
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Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar,
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/clone_lib="UT0064"
/dev_stage="Adult"
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GGGACCAATGATGGTGCACAAAGGCTTAGATTTGCCTTGTCTCAAAATAAGGAATTTTGT
                                                                                                cDNA Library Preparation: Life Technologies, Inc cDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AW189584 538 bp mRNA EST 22-NOV-1999 x108h12.x1 NCI_CGAP_Ut4 Homo sapiens cDNA clone IMAGE:2675687 3'
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Seq primer: -40UP from Gibco
High quality sequence stop: 175.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tissue Procurement: Christopher Moskaluk, M.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (1997)
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National Cancer Institute, Cancer Genome Anai
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D.,
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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clarity 99.8%;
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Site_2: Notl; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.69 kb. Life Technologies catalog *:
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/clone="IMAGE:2623947"
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/tlssue_type-"poorly differentiated adenocarcinoma with
                                                                                                                                                                                                                                                                                    /lab_host-"DH10B"
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                                                                                                                                                                            Possible reversed clone: polyT not found Seq primer: -40UP from Gibco
                                                                                                                                                                                                                                                                                                                                   Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D.,
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 475)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AW589567 475 bp mRNA EST 22-MAR-2000 xol2bl2.xl NCI_CGAP_Ut3 Homo sapiens cDNA clone IMAGE:2703743 similar to TR:095786 095786 RNA HELICASE. ;, mRNA sequence.
                                                                                                                                                          High quality sequence stop: 407.
                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                               Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                           Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                              National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCI-CGAP http://www.ncbi.nlm.nih.gov/nciegap
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cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
                                                                                                                                                                                                                                                                                                                                                                                                                              Gene Index
/clone="IMAGE:2703743"
/clone_lib="NCI_CGAP_Ut3"
/tissue_type="poorly-differentiated endometrial adenocarcinoma, 2 pooled tumors"
/lab_host="DH10B"
                                                                                       /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                        Location/Qualifiers
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                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 479)
                                                                                                                     Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Chris Moskaluk, M.D., Ph.D., Michael
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                Tumor Gene Index
Unpublished (1997)
                                                                                                                                                                            Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                            Homo sapiens
                  info@image.llnl.gov
                             found through the I.M.A.G.E. Consortium/LLNL,
                                                CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
                                                                                                                                                                                                                                    National Cancer Institute, Cancer
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-40UP from Gibco
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100.0%;
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                                   send email to:
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atttcccaatcttgactattcagaatgctgtttatttagtgatgaggattagcacttgat 3254
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                                                                                              tgaagattettttaaaataeetateagttaaacatttaatatgattatgattaatgtatte
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470; Conserv
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /db_xref="taxon:9666"
/dlone="IMAGE:364554"
/clone=!ib="MCI_GAP_BF22"
/clone_lib="MCI_GAP_BF22"
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/note="forgan: breast; Vector: pCMV-SPORT6; Site_1: Not1;
Site_2: Sal1; Cloned unidirectionally. Primer: Oligo dT.
average insert size 1.5 kb. Constructed by Life
Technologies."
71 g 179 t
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Pred. No. 3.5e-86;
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Search completed: January 17, 2002, 02:29:15 Job time: 4399 sec · ·